

AMENDMENTS TO THE CLAIMS:

Please replace the claims with the claims provided in the listing below wherein status, amendments, additions and cancellations are indicated.

1. (Currently amended) ~~A method~~ Method for recording sequence information on biological compounds, comprising the steps of:

dividing [[the]] one of text data representing [[the]] a sequence of said biological compounds ~~or the~~ and numerical data obtained by converting said text data ~~based on, utilizing~~ a conversion rule, into a plurality of m-bit partial data arranged in a plurality of columns in [[the]] an arranged direction corresponding to [[the]] a direction along which said biological compounds are placed and in a plurality of rows in [[the]] a non-arranged direction which crosses said arranged direction, where m is an integer larger greater than or equal to 16;

computing a first set of parity information by applying a first operation of a Galois field $GF(2^m)$ along said non-arranged direction to a set of said partial data of each column;

computing a second set of parity information by applying a second operation of a Galois field $GF(2^m)$ along said arranged direction to a set of said partial data of each row; and

representing said sequence information on said biological compounds by

said first and second sets of parity information.

2. (Currently amended) ~~The method~~ Method of claim 1, wherein
- ~~assuming that when~~ α is a primitive element of a Galois field GF(2^m),
- said first set of parity information includes [[the]] a sum of a plurality of products obtained by multiplying a set of said partial data of each column along said non-arranged direction by α^{sp} , $\alpha^{s(p+1)}$, $\alpha^{s(p+2)}$, ..., $\alpha^{s(p+dp)}$, where s and p are nonnegative integers and dp is an integer ~~lager~~ greater than or equal to one; and
- said second set of parity information includes [[the]] a sum of a plurality of products obtained by multiplying a set of said partial data of each row along said arranged direction by α^{tq} , $\alpha^{t(q+1)}$, $\alpha^{t(q+2)}$, ..., $\alpha^{t(q+dq)}$, where t and q are nonnegative integers and dq is an integer ~~lager~~ greater than or equal to one.

3. (Currently amended) ~~The method~~ Method of claim 2, wherein
- [[the]] a number of said partial data placed along said arranged direction is smaller than [[the]] a number of said partial data placed along said non-arranged direction; and
- [[the]] a number of said second set of parity information is smaller than [[the]] a number of said first set of parity information.

4. (Currently amended) ~~The method~~ Method of claim 3, wherein [[the]] said number of said partial data placed along said non-arranged direction is smaller less than or equal to $(2^m-1)/4$.

5. (Currently amended) ~~The method~~ Method of claim 2, wherein both of said integers s and t are zero.

6. (Currently amended) ~~The method~~ Method of claim 2, wherein both of said integers s and t are one.

7. (Currently amended) ~~The method~~ Method of claim 2, wherein said first set of parity information includes a plurality of said sums obtained for each column using mutually different values of said integer s; and said second set of parity information includes a plurality of said sums obtained for each row using mutually different values of said integer t.

8. (Currently amended) ~~The method~~ Method of claim 1, wherein said partial data is said numerical data obtained by expressing each of said biological compounds [[in]] as data whose having a size [[is]] less than or equal to six bits.

9. (Currently amended) ~~The method~~ Method of claim 1, wherein said integer m that defines said Galois field GF(2^m) is a multiple of 64.

10. (Currently amended) ~~The method~~ Method of claim 1, further including comprising the steps of:

assuming making an assumption that said sequence of said biological compounds is a standard sequence;

computing two sets of parity information on a sequence of biological compounds subject to examination correspondingly, said two sets of parity information on said sequence of biological compounds subject to examination corresponding to said two sets of parity information on said standard sequence; and

identifying [[the]] differences between said standard sequence and said sequence of biological compounds subject to examination, by using said [[four]] two sets of parity information on said standard sequence and said two sets of parity information on said sequence of biological compounds subject to examination.

11. (Currently amended) ~~The method~~ Method of claim 1, wherein said biological compounds are nucleotides constituting at least part of DNA, RNA, or a gene.

12. (Currently amended) ~~The method~~ Method of claim 1, wherein said biological compounds are amino acids constituting at least part of a protein.

13. (Currently amended) ~~A device~~ Device for recording sequence information on biological compounds, comprising:

a sequencer for reading sequence information on said biological compounds;

dividing means for dividing [[the]] one of text data representing [[the]] a sequence of said biological compounds ~~or the~~ and numerical data obtained by converting said text data ~~based on~~ utilizing a conversion rule, into a plurality of m-bit partial data arranged in a plurality of columns in [[the]] an arranged direction corresponding to [[the]] a direction along which said biological compounds are placed, and in a plurality of rows in [[the]] a non-arranged direction which crosses said arranged direction, where m is an integer larger greater than or equal to 16;

computing means for computing a first set of parity information by applying a first operation of a Galois field GF(2^m) along said non-arranged direction to a set of said partial data of each column and computing a second set of parity information by applying a second operation of a Galois field GF(2^m) along said arranged direction to a set of said partial data of each row; and

recording means for recording said first and second sets of parity information in a recording medium.

14. (Currently amended) ~~The device~~ Device of claim 13, wherein
assuming that when α is a primitive element of a Galois field $GF(2^m)$,
said first set of parity information includes [[the]] a sum of a plurality of products obtained by multiplying a set of said partial data of each column along said non-arranged direction by α^{sp} , $\alpha^{s(p+1)}$, $\alpha^{s(p+2)}$, ..., $\alpha^{s(p+dp)}$, where s and p are nonnegative integers and dp is an integer lager greater than or equal to one; and
said second set of parity information includes [[the]] a sum of a plurality of products obtained by multiplying a set of said partial data of each row along said arranged direction by α^{tq} , $\alpha^{t(q+1)}$, $\alpha^{t(q+2)}$, ..., $\alpha^{t(q+dq)}$, where t and q are nonnegative integers and dq is an integer lager greater than or equal to one.

15. (Currently amended) ~~A computer-readable~~ Computer-readable medium ~~for~~ storing sequence information on biological compounds, comprising:

a data structure stored in said medium, wherein ~~in order to form said data structure,~~
[[the]] one of text data representing [[the]] a sequence of said biological compounds ~~or the and~~ numerical data obtained by converting said text data ~~based~~

on utilizing a conversion rule is divided into a plurality of m-bit partial data arranged in a plurality of columns in [[the]] an arranged direction, corresponding to [[the]] a direction along which said biological compounds are placed, and in a plurality of rows in [[the]] a non-arranged direction which crosses said arranged direction, where m is an integer larger greater than or equal to 16;

a first set of parity information is computed by applying a first operation of a Galois field GF(2^m) along said non-arranged direction to a set of said partial data of each column and a second set of parity information is computed by applying a second operation of a Galois field GF(2^m) along said arranged direction to a set of said partial data of each row; and

said first and second sets of parity information are recorded in said data structure as said sequence information on said biological compounds.

16. (Currently amended) The computer-readable Computer-readable medium of claim 15, wherein

said data structure further includes a mathematical digest of one of said text data representing said sequence of said biological compounds [[or]] and said numerical data corresponding to said text data, where and

~~the size of~~ said mathematical digest is larger has a size greater than or equal to 40 bits.

17. (Currently amended) A method Method for supplying sequence information on biological compounds, comprising the steps of:

~~as the procedure of a supplier;~~

recording [[the]] one of text data representing [[the]] a sequence of said biological compounds ~~or the and~~ numerical data obtained by converting said text data ~~based on utilizing~~ a conversion rule, in a first file;

dividing said one of text data and numerical data recorded in said first file ~~or said numerical data recorded in said first file~~ into a plurality of m-bit partial data arranged in a plurality of columns in [[the]] an arranged direction corresponding to [[the]] a direction along which said biological compounds are placed, and in a plurality of rows in [[the]] a non-arranged direction which crosses said arranged direction, where m is an integer larger greater than or equal to 16;

computing a first set of parity information by applying a first operation of a Galois field $GF(2^m)$ along said non-arranged direction to a set of said partial data of each column, and computing a second set of parity information by applying a second operation of a Galois field $GF(2^m)$ along said arranged direction to a set of said partial data of each row;

recording said first and second sets of parity information in a second file;
and

~~as the procedure of a user,~~
~~receiving providing~~ said two sets of parity information recorded in said second file through a communications network ~~from said supplier.~~

18. (Currently amended) ~~The method of claim 17, further including Method for utilizing sequence information on biological compounds, comprising the steps of:~~

~~as the procedure of said user, receiving through said communications network from a supplier~~ said two sets of parity information recorded on said second file, according to claims 17;

identifying [[the]] differences between said sequence of said biological compounds held by said supplier and [[the]] a sequence of biological compounds subject to examination, based on said two sets of received parity information; and

when said differences cannot be recovered, ~~said user~~ receiving [[the]] sequence information on [[the]] a part corresponding to said differences, within said one of text data and numerical data recorded in said first file or ~~said numerical data recorded in said first file~~, through said communications network from said supplier.

19. (Currently amended) ~~The method~~ Method of claim 18, wherein

assuming that when α is a primitive element of a Galois field GF(2^m), said first set of parity information includes [[the]] a sum of a plurality of products obtained by multiplying a set of said partial data of each column along said non-arranged direction by α^{sp} , $\alpha^{s(p+1)}$, $\alpha^{s(p+2)}$, ..., $\alpha^{s(p+dp)}$, where s and p are nonnegative integers and dp is an integer larger greater than or equal to one; and said second set of parity information includes [[the]] a sum of a plurality of products obtained by multiplying a set of said partial data of each row along said arranged direction by α^{tq} , $\alpha^{t(q+1)}$, $\alpha^{t(q+2)}$, ..., $\alpha^{t(q+dq)}$, where t and q are nonnegative integers and dq is an integer larger greater than or equal to one.

20. (Currently amended) The method Method of claim [[18]] 17, further including comprising the steps step of:

as the procedure of said supplier,
letting the providing information on [[the]] a number of said sequence of said biological compounds, and [[the]] information on a mathematical digest of said one of text data or said and numerical data be disclosed to the public through said communications network; and

~~as the procedure of a user,~~
~~accessing said information on the number of said sequence of biological~~
~~compounds and said information on said mathematical digest through said communications~~
~~network before receiving said two sets of parity information.~~

21. (Currently amended) ~~A method~~ Method for recording sequence information on biological compounds, comprising the steps of:

dividing [[the]] one of text data representing [[the]] a sequence of said biological compounds ~~or the and~~ numerical data obtained by converting said text data ~~based on utilizing~~ a conversion rule_s into a plurality of m-bit partial data arranged in a plurality of columns in [[the]] an arranged direction corresponding to [[the]] a direction along which said biological compounds are placed_s and in a plurality of rows in [[the]] a non-arranged direction which crosses said arranged direction, where m is an integer larger greater than or equal to 16;

assuming making assumptions that a maximum value of said partial data is Nmax_s and a prime number larger than said maximum value Nmax is P;

computing a first set of parity information by applying a first operation of a Galois field GF(P) along said non-arranged direction to a set of said partial data of each column;

computing a second set of parity information by applying a second operation of a Galois field GF(P) along said arranged direction to a set of said partial data of each row; and

representing said sequence information on said biological compounds by said first and second sets of parity information.

22. (Currently amended) ~~The method Method~~ of claim 21, wherein
said maximum value Nmax of said partial data is smaller than ($2^m - 1$) and
said prime number P satisfies the following condition:

$$2^m > P > Nmax.$$

23. (Currently amended) ~~The method Method~~ of claim 22, wherein
~~assuming that when~~ δ is a primitive element of a Galois field GF(P),
said first set of parity information includes [[the]] a sum of a plurality of
products obtained by multiplying a set of said partial data of each column along
said non-arranged direction by δ^{sp} , $\delta^{s(p+1)}$, $\delta^{s(p+2)}$, ..., $\delta^{s(p+dp)}$, where s and p are
nonnegative integers and dp is an integer ~~lager~~ greater than or equal to one; and
said second set of parity information includes [[the]] a sum of a plurality
of products obtained by multiplying a set of said partial data of each row along said
arranged direction by δ^{tq} , $\delta^{t(q+1)}$, $\delta^{t(q+2)}$, ..., $\delta^{t(q+dq)}$, where t and q are nonnegative
integers and dq is an integer ~~lager~~ greater than or equal to one.

24. (Currently amended) ~~A method Method~~ for supplying sequence
information on biological compounds, comprising the steps of:
~~as the procedure of a supplier;~~
recording [[the]] one of text data representing the sequence of said

biological compounds ~~or the~~ and numerical data obtained by converting said text data ~~based on~~ utilizing a conversion rule, in a first file;

dividing said one of said text data and said numerical data recorded in said first file ~~or said numerical data recorded in said first file~~ into a plurality of m-bit partial data arranged in a plurality of columns in [[the]] an arranged direction corresponding to [[the]] a direction along which said biological compounds are placed, and in a plurality of rows in [[the]] a non-arranged direction which crosses said arranged direction, where m is an integer larger than or equal to 16;

assuming making assumptions that a maximum value of said partial data is Nmax, and a prime number larger than said maximum value Nmax is P;

computing a first set of parity information by applying a first operation of a Galois field GF(P) along said non-arranged direction to a set of said partial data of each column and computing a second set of parity information by applying a second operation of a Galois field GF(P) along said arranged direction to a set of said partial data of each row;

recording said first and second sets of parity information in a second file; and

~~as the procedure of a user,~~

~~receiving providing~~ said two sets of parity information recorded in said second file through a communications network ~~from said supplier.~~

25. (Currently amended) ~~The method of claim 24, further including Method for utilizing sequence information on biological compounds comprising the steps of:~~

~~as the procedure of said user, receiving through said communications network from a supplier said two sets of parity information recorded in said second file, according to claim 24;~~

identifying [[the]] differences between said sequence of said biological compounds held by said supplier and [[the]] a sequence of biological compounds subject to examination, based on said two sets of received parity information and

when said differences cannot be recovered, ~~said user~~ receiving [[the]] sequence information on [[the]] a part corresponding to said differences, within said one of text data recorded in said first file ~~or said numerical data recorded in said first file,~~ through said communications network from said supplier

26. (New) Method of claim 18, further comprising the step of:

accessing at least one of information on a number of said sequence of said biological compounds, and information on a mathematical digest of said one of text data and numerical data, through said communications network.

CLAIMS

What is claimed is:

- 5 1. A method for recording sequence information on biological compounds, comprising the steps of:
 - dividing the text data representing the sequence of said biological compounds or the numerical data obtained by converting said text data based on a conversion rule into a plurality of m-bit partial data arranged in a plurality of columns in the arranged direction corresponding to the direction along which said biological compounds are placed and in a plurality of rows in the non-arranged direction which crosses said arranged direction, where m is an integer larger than or equal to 16;
 - 10 computing a first set of parity information by applying a first operation of Galois field GF(2^m) along said non-arranged direction to a set of said partial data of each column;
 - 15 computing a second set of parity information by applying a second operation of Galois field GF(2^m) along said arranged direction to a set of said partial data of each row; and
 - 20 representing said sequence information on said biological compounds by said first and second sets of parity information.
2. The method of claim 1, wherein
 - assuming that α is a primitive element of Galois field GF(2^m),
 - 25 said first set of parity information includes the sum of a plurality of products obtained by multiplying a set of said partial data of each column along said non-arranged direction by α^{sp} , $\alpha^{s(p+1)}$, $\alpha^{s(p+2)}$, ..., $\alpha^{s(p+dp)}$, where s and p are nonnegative integers and dp is an integer lager than or equal to one; and
 - 30 said second set of parity information includes the sum of a plurality of products obtained by multiplying a set of said partial data of each row along said arranged direction by α^{tq} , $\alpha^{t(q+1)}$, $\alpha^{t(q+2)}$, ..., $\alpha^{t(q+dq)}$, where t and q are nonnegative integers and dq is an integer lager than or equal to one.
3. The method of claim 2, wherein

the number of said partial data placed along said arranged direction is smaller than the number of said partial data placed along said non-arranged direction; and

5 the number of said second set of parity information is smaller than the number of said first set of parity information.

4. The method of claim 3, wherein the number of said partial data placed along said non-arranged direction is smaller than or equal to $(2^m \cdot 1)/4$.

10 5. The method of claim 2, wherein both of said integers s and t are zero.

6. The method of claim 2, wherein both of said integers s and t are one.

7. The method of claim 2, wherein

15 said first set of parity information includes a plurality of said sums obtained for each column using mutually different values of said integer s; and

said second set of parity information includes a plurality of said sums obtained for each row using mutually different values of said integer t.

20 8. The method of claim 1, wherein said partial data is said numerical data obtained by expressing each of said biological compounds in data whose size is less than or equal to six bits.

25 9. The method of claim 1, wherein said integer m that defines Galois field GF(2^m) is a multiple of 64.

10. The method of claim 1, further including the steps of:

assuming that said sequence of said biological compounds is a standard sequence;

30 computing two sets of parity information on a sequence of biological compounds subject to examination correspondingly to said two sets of parity information on said standard sequence; and

identifying the differences between said standard sequence and said sequence of biological compounds subject to examination by using said four sets of

parity information.

11. The method of claim 1, wherein said biological compounds are nucleotides constituting at least part of DNA, RNA, or a gene.

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12. The method of claim 1, wherein said biological compounds are amino acids constituting at least part of a protein.

13. A device for recording sequence information on biological compounds,
10 comprising:

a sequencer for reading sequence information on said biological compounds;

dividing means for dividing the text data representing the sequence of said biological compounds or the numerical data obtained by converting said text data based on a conversion rule into a plurality of m-bit partial data arranged in a plurality of columns in the arranged direction corresponding to the direction along which said biological compounds are placed and in a plurality of rows in the non-arranged direction which crosses said arranged direction, where m is an integer larger than or equal to 16;

20 computing means for computing a first set of parity information by applying a first operation of Galois field $GF(2^m)$ along said non-arranged direction to a set of said partial data of each column and computing a second set of parity information by applying a second operation of Galois field $GF(2^m)$ along said arranged direction to a set of said partial data of each row; and

25 recording means for recording said first and second sets of parity information in a recording medium.

14. The device of claim 13, wherein

assuming that α is a primitive element of Galois field $GF(2^m)$,

30 said first set of parity information includes the sum of a plurality of products obtained by multiplying a set of said partial data of each column along said non-arranged direction by α^{sp} , $\alpha^{s(p+1)}$, $\alpha^{s(p+2)}$, ..., $\alpha^{s(p+dp)}$, where s and p are nonnegative integers and dp is an integer larger than or equal to one; and

said second set of parity information includes the sum of a plurality of

products obtained by multiplying a set of said partial data of each row along said arranged direction by α^{tq} , $\alpha^{t(q+1)}$, $\alpha^{t(q+2)}$, ..., $\alpha^{t(q+dq)}$, where t and q are nonnegative integers and dq is an integer larger than or equal to one.

5 15. A computer-readable medium storing sequence information on biological compounds, comprising:

a data structure stored in said medium, wherein in order to form said data structure,

10 the text data representing the sequence of said biological compounds or the numerical data obtained by converting said text data based on a conversion rule is divided into a plurality of m-bit partial data arranged in a plurality of columns in the arranged direction corresponding to the direction along which said biological compounds are placed and in a plurality of rows in the non-arranged direction which crosses said arranged direction, where m is an integer larger than or equal to 16;

15 a first set of parity information is computed by applying a first operation of Galois field GF(2^m) along said non-arranged direction to a set of said partial data of each column and a second set of parity information is computed by applying a second operation of Galois field GF(2^m) along said arranged direction to a set of said partial data of each row; and

20 said first and second sets of parity information are recorded in said data structure as said sequence information on said biological compounds.

16. The computer-readable medium of claim 15, wherein

25 said data structure further includes a mathematical digest of said text data representing said sequence of said biological compounds or said numerical data corresponding to said text data, where

the size of said mathematical digest is larger than or equal to 40 bits.

30 17. A method for supplying sequence information on biological compounds, comprising the steps of:

as the procedure of a supplier,

recording the text data representing the sequence of said biological compounds or the numerical data obtained by converting said text data based on a conversion rule in a first file;

dividing said text data recorded in said first file or said numerical data recorded in said first file into a plurality of m-bit partial data arranged in a plurality of columns in the arranged direction corresponding to the direction along which said biological compounds are placed and in a plurality of rows in the non-arranged direction which crosses said arranged direction, where m is an integer larger than or equal to 16;

computing a first set of parity information by applying a first operation of Galois field GF(2^m) along said non-arranged direction to a set of said partial data of each column and computing a second set of parity information by applying a second operation of Galois field GF(2^m) along said arranged direction to a set of said partial data of each row;

recording said first and second sets of parity information in a second file; and

as the procedure of a user,

receiving said two sets of parity information recorded in said second file through a communications network from said supplier.

18. The method of claim 17, further including the steps of:

as the procedure of said user,

identifying the differences between said sequence of said biological compounds held by said supplier and the sequence of biological compounds subject to examination based on said two sets of parity information; and

when said differences cannot be recovered, said user receiving the sequence information on the part corresponding to said differences within said text data recorded in said first file or said numerical data recorded in said first file through said communications network from said supplier.

19. The method of claim 18, wherein

assuming that α is a primitive element of Galois field GF(2^m),

said first set of parity information includes the sum of a plurality of products obtained by multiplying a set of said partial data of each column along said non-arranged direction by α^{sp} , $\alpha^{s(p+1)}$, $\alpha^{s(p+2)}$, ..., $\alpha^{s(p+dp)}$, where s and p are nonnegative integers and dp is an integer larger than or equal to one; and

said second set of parity information includes the sum of a plurality of

products obtained by multiplying a set of said partial data of each row along said arranged direction by α^{tq} , $\alpha^{t(q+1)}$, $\alpha^{t(q+2)}$, ..., $\alpha^{t(q+dq)}$, where t and q are nonnegative integers and dq is an integer larger than or equal to one.

- 5 20. The method of claim 18, further including the steps of:
 as the procedure of said supplier,
 letting the information on the number of said sequence of said biological
 compounds and the information on a mathematical digest of said text data or said
 numerical data be disclosed to the public through said communications network; and
10 as the procedure of a user,
 accessing said information on the number of said sequence of biological
 compounds and said information on said mathematical digest through said
 communications network before receiving said two sets of parity information.
- 15 21. A method for recording sequence information on biological compounds,
 comprising the steps of:
 dividing the text data representing the sequence of said biological
 compounds or the numerical data obtained by converting said text data based on a
 conversion rule into a plurality of m-bit partial data arranged in a plurality of
20 columns in the arranged direction corresponding to the direction along which said
 biological compounds are placed and in a plurality of rows in the non-arranged
 direction which crosses said arranged direction, where m is an integer larger than or
 equal to 16;
 assuming that a maximum value of said partial data is Nmax
25 and a prime number larger than said maximum value Nmax is P;
 computing a first set of parity information by applying a first
 operation of Galois field GF(P) along said non-arranged direction to a set of said
 partial data of each column;
 computing a second set of parity information by applying a
30 second operation of Galois field GF(P) along said arranged direction to a set of said
 partial data of each row; and
 representing said sequence information on said biological
 compounds by said first and second sets of parity information.

22. The method of claim 21, wherein

said maximum value Nmax of said partial data is smaller than $(2^m - 1)$ and said prime number P satisfies the following condition:

$$2^m > P > Nmax.$$

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23. The method of claim 22, wherein

assuming that δ is a primitive element of Galois field GF(P),

10 said first set of parity information includes the sum of a plurality of products obtained by multiplying a set of said partial data of each column along said non-arranged direction by δ^{sp} , $\delta^{s(p+1)}$, $\delta^{s(p+2)}$, ..., $\delta^{s(p+dp)}$, where s and p are nonnegative integers and dp is an integer larger than or equal to one; and

15 said second set of parity information includes the sum of a plurality of products obtained by multiplying a set of said partial data of each row along said arranged direction by δ^{tq} , $\delta^{t(q+1)}$, $\delta^{t(q+2)}$, ..., $\delta^{t(q+dq)}$, where t and q are nonnegative integers and dq is an integer larger than or equal to one.

24. A method for supplying sequence information on biological compounds, comprising the steps of:

as the procedure of a supplier,

20 recording the text data representing the sequence of said biological compounds or the numerical data obtained by converting said text data based on a conversion rule in a first file;

25 dividing said text data recorded in said first file or said numerical data recorded in said first file into a plurality of m-bit partial data arranged in a plurality of columns in the arranged direction corresponding to the direction along which said biological compounds are placed and in a plurality of rows in the non-arranged direction which crosses said arranged direction, where m is an integer larger than or equal to 16;

30 assuming that a maximum value of said partial data is Nmax and a prime number larger than said maximum value Nmax is P;

computing a first set of parity information by applying a first operation of Galois field GF(P) along said non-arranged direction to a set of said partial data of each column and computing a second set of parity information by applying a second operation of Galois field GF(P) along said arranged direction to a set of said partial

data of each row;

recording said first and second sets of parity information in a second file; and

as the procedure of a user,

5 receiving said two sets of parity information recorded in said second file through a communications network from said supplier.

25. The method of claim 24, further including the steps of:

as the procedure of said user,

10 identifying the differences between said sequence of said biological compounds held by said supplier and the sequence of biological compounds subject to examination based on said two sets of parity information; and

when said differences cannot be recovered, said user receiving the sequence information on the part corresponding to said differences within said text data 15 recorded in said first file or said numerical data recorded in said first file through said communications network from said supplier.